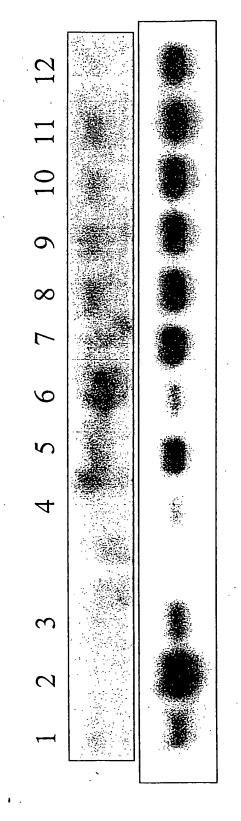
mRNA Reverse Transcriptase Primers: YT.....<sub>15</sub>U<sub>2</sub>AG,  $G_7AU_2dT$ : cDNA • (cDNA normalization by PCR with Primers: CAPDH  $G_7AU_2T$ :  $G_7TATTTATTTAT$  $T_{15}U_2AG_n$ Final product 6% native PAAG Fragment elution, PCR, cloning and verification of differential expression Sequence analysis, cDNA cloning





2. Skeletal muscle 4. Spleen 3. Liver

1. Colon

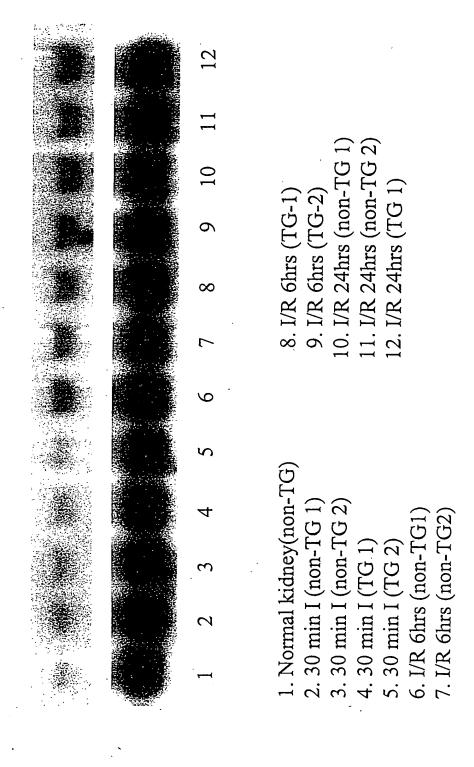
5. Small Intestine

6. Ovary 7. Brain 8-12. Kidney samples

Lower panel: the same blot hybridized with mouse GAPDH gene probe, shown as Upper panel: Northern blot hybridized with candidate gene probe internal control

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Figure 3



Express Mail No. EL740476596US Docket No. 266/171 Sheet No. 3 of 7 1. Molecular marker

Total cell extract

20,000g ppt after sonication

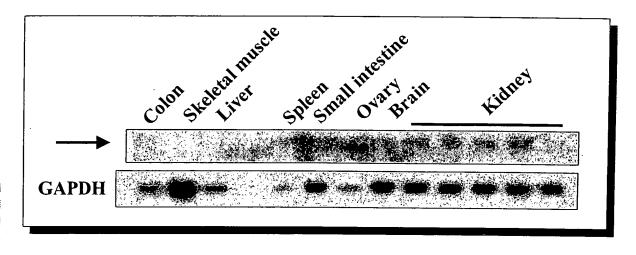
180,000g ppt

Below States

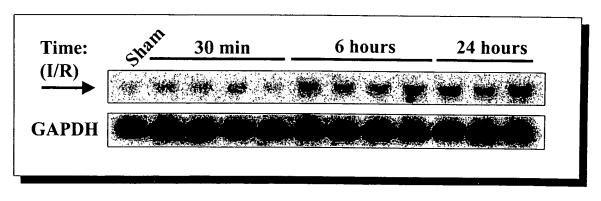
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## mRNA expression in mouse tissues

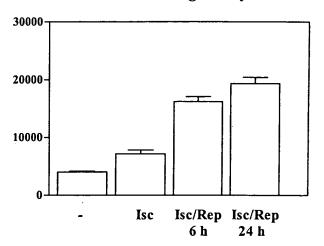


## mRNA expression after kidney ischemia/reperfusion



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## Measurement of IAP mRNA activation during kidney I/R \*

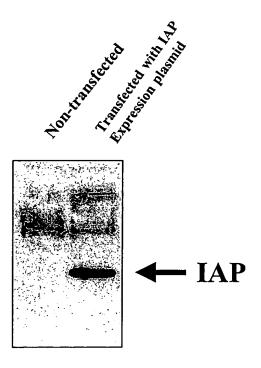


\*At least 4 animals were analyzed per each group.

Data are expressed as the mean +SEM.

Figure 6

## Overexpression of IAP in Hela cells



Hela cells were transfected with plasmid DNA, containing pEF/myc/cyto vector (Invitrogen) containing human IAP cDNA. After 48 hrs cells were lysed and total cellular extract was analyzed by Western blotting using anti-IAP rabbit polyclonal Ab and Phototope-HRP Western Blot Detection Kit (BioLabs) as described in manufacturer's protocol.